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GENENTECH, INC. LEGAL DEPT.





5 DEPARTMENT OF COMMERCE Patent and Trademark Office

COMMISSIONER OF PATENTS AND TRADEMARKS

Washington, D.C. 20231

ATTORNEY DOCKET NO.

09/313,299

05/17/99

L.F.F.

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APPLICATION NO.

HMCD/FLVW

FIRST NAMED INVENTOR

计算机 经现代证券 计

TIMOTHY R SCHWORTS FH D

GENERICECH IM.

1 DMR WAY

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EXAMINER MUTCHER **ART UNIT** PAPER NUMBER 16016

DATE MAILED:

West Day Kills

Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner of Patents and Trademarks

CALENDARED/8



UNITED COMMERCE Patent and Trademark Office COMMISSIONER OF PATENTS AND TRADEMARKS Washington, D.C. 20231

09313299 SERIAL NUMBER FILING DATE	FIRST NAMED APPLICANT	ATTORNEY DOCKET NO.
Γ .	1	EXAMINER
	ARTIIN	T PAPER NUMBER

DATE MAILED:

ART UNIT

Please find below a communication from the EXAMINER in charge of this application

The communication filed on 11/22/99 is not fully responsive to the communication mailed 10/29/99 for the reason(s) set forth on the attached Notice to Comply With the Sequence Rules or CRF Diskette Problem Report.

Since the response appears to be **bona** fide, but through an apparent oversight or inadvertence failed to provide a complete response, applicant is given ONE (1) MONTH from the mailing date of this notice within which to supply the omission or correction in order to avoid abandonment. EXTENSIONS OF THIS TIME PERIOD MAY BE GRANTED UNDER 37 CFR 1.136(a).

Any inquiry concerning this communication should be directed to Examiner Joseph F. Murphy, Art Unit 1644, whose telephone number is 703-305-7245.

Any inquiry of a general nature or relating to the status of this application should be directed to the Technology Center receptionist whose telephone number is (703) 308-0196.

Joseph F. Murphy, Ph.D.

Patent Examiner Art Unit 1644

L

December 30, 1999

PRIMARY EXAMINER





GAU 1644 WD

Patent Docket No. P0963R1D1

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of

James Lee et al.

Serial No.: 09/313,299

Filed: May 17, 1999

For: VEGF - RELATED PROTEIN

Examiner: J. Murphy

Art Unit: 1644

CERTIFICATE OF MAILING

I hereby certify that this correspondence is being deposited with the United States Postal Service with sufficient postage as first class mail in an envelope addressed to: Assistant Commissioner for Patents, Washington, DC 20231 on

March / 3, 2000

Tara E. Faenza

TRANSMITTAL LETTER

BOX SEQUENCE
Assistant Commissioner for Patents
Washington, DC 20231

MAR 2 3 2000

TECH CENTER 1600/2900

Sir:

Enclosed for filing in the above-identified patent application, please find:

- 1. Response to Notice to Comply with the Requirement for Patent Application Containing Sequence Disclosures;
- 2. Copy of Notice to Comply with the Requirement for Patent Application Containing Sequence Disclosures;
 - 3. Sequence Listing and computer-readable diskette;
 - 4. Petition for One Month Extension of Time.
 - 5. A return postage-prepaid postcard.

The Commissioner is authorized to charge any additional fees due in connection with the filing of these documents to our Deposit Account No. 07-0630 (our Docket P0963R1D1). <u>A duplicate copy of this transmittal is enclosed</u>.

By:_

Respectfully submitted,

GENENTECH, INC.

Date: March (3, 2000)

Timothy R. Schwartz, Ph.D.

Reg. No. 32,171

1 DNA Way

South San Francisco, CA 94080-4990

Phone: (650) 225-7467 Fax: (650) 952-9881 MAR 2 1 2000

TECH CENTER 1600/290



Patent Docket No. P0963R1D1

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of

James Lee et al.

Serial No.: 09/313,299

Filed: May 17, 1999

For: VEGF - RELATED PROTEIN

Examiner: J. Murphy

Art Unit: 1644

CERTIFICATE OF M'AILING

I hereby certify that this correspondence is being deposited with the United States Postal Service with sufficient postage as first class mail in an envelope addressed to: Assistant Commissioner for Patents, Washington, DC 20231 on

March 12, 2000

RESPONSE TO NOTICE TO COMPLY WITH REQUIREMENTS FOR

PATENT APPLICATION CONTAINING SEQUENCE DISCLOSURES

BOX SEQUENCE Assistant Commissioner for Patents Washington, DC 20231

Sir:

In response to the Notice to Comply with Requirements for Patent Applications Containing Nucleotiude Sequence and/or Amino Acid Sequence Disclosures mailed January 24, 2000, transmitted herewith are the following documents:

- 1. Copy of Notice to Comply with Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures;
 - 2. Sequence Listing and computer-readable diskette;

The Commissioner is authorized to charge any additional fees due in connection with the filing of these documents to our Deposit Account No. 07-0630 (our Docket P0963R1D1). <u>A duplicate copy of this transmittal is enclosed</u>.

Respectfully submitted,

GENENTECH, INC.

Date: March 12, 2000

Timothy R. Schwartz, Ph.D

Reg. No. 32,171

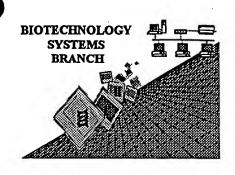
1 DNA Way

South San Francisco, CA 94080-4990

Phone: (650) 225-7467 Fax: (650) 952-9881

#72020

RAW SEQUENCE LISTING O I P ERROR REPORT MAR 2 0 2000 E



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/313,299A

Art Unit / Team No. :

1646

Date Processed by STIC:

12/10/99

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

Raw Sequence Listing Error Summary

	ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 09/3/3, 299/
ATTN.	NEW RILLES CASES: P	LEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
1	Wrapped Nucleics	The number/text at the end of each line "wrapped" down to the next line.
·		This may occur if your file was retrieved in a word processor after creating it.
		Please adjust your right margin to .3, as this will prevent "wrapping".
2	Wrapped Aminos	The amino acid number/text at the end of each line "wrapped " down to the next line:
• ——		This may occur if your file was retrieved in a word processor after creating it.
		Please adjust your right margin to .3, as this will prevent "wrapping".
3	Incorrect Line Length	The rules require that a line not exceed 72 characters in length. This includes spaces.
4	Misaligned Amino Acid	The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs
'	Numbering	between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
5	Non-ASCII	This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
		Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
6	Variable Length	Sequence(s) contain n's or Xaa's which represented more than one residue.
	-	As per the rules, each n or Xaa can only represent a single residue.
		Please present the maximum number of each residue having variable length and
		indicate in the (ix) feature section that some may be missing.
7	Patentin ver. 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
		sequence(s) Normally, Patentin would automatically generate this section from the
		previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section
		to the subsequent amino acid sequence.
8	Skipped Sequences	Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
	(OLD RULES)	(2) INFORMATION FOR SEQ ID NO:X:
		(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
		(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
		This sequence is intentionally skipped
		Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
9	Skipped Sequences	Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
	(NEW RULES)	<210> sequence id number
		<400> sequence id number
		000
0	Use of n's or Xaa's	Use of n's and/or Xaa's have been detected in the Sequence Listing.
	(NEW RULES)	Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
		In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
1	Use of <213>Organism	Sequence(s) are missing this mandatory field or its response.
1	(NEW RULES)	1-9 (maybe more)
2	Use of <220>Feature	Sequence(s) are missing the <220>Feature and associated headings.
	(NEW RULES)	Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
		Please explain source of genetic material in <220> to <223> section.
		(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
3	Patentin ver. 2.0 "bug"	Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted
		file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
		Instead, please use "File Manager" or any other means to copy file to floppy disk.

AKS-Biotechnology Systems Branch- 5/15/99

Application No.: <u>69313255</u>

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

E].	 This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990. 		
		2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).		
]	3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).		
Þ	1	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked-up "Raw Sequence Listing."		
]	The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).		
		The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).		
		7. Other:		
A	Applicant Must Provide:			
	$\overline{\mathbb{Z}}$	An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".		
[An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.		
1	3	A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).		

For questions regarding compliance to these requirements, please contact

For Rules Interpretation, call (703) 308-4216

For CRF Submission Help, call (703) 308-4212

For Patentin software help, call (703) 308-6856

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE